Truncated distributions

35 marks

The questions here are designed to explore some basic characteristics of, and differences between, probability distributions and the random realizations from them. See help("Distributions") for those built into R.

1. Suppose we have a continuous random variable X with distribution function $F_X(x) = Pr(X \le x)$ and quantile function $Q_X(p) = F_X^{-1}(p)$. That is $p = F_X(x) = Pr(X \le x)$ and $p = Pr(X \le Q_X(p)) = F_X(Q_X(p)) = F_X(F_X^{-1}(p)) = p$.

We can define a random variable Y having cumulative distribution function

$$G_Y(y) = \begin{cases} 0 & y < a \\ \frac{F_X(y) - F_X(a)}{F_X(b) - F_X(a)} & a \le y \le b \\ 1 & y > b \end{cases}$$

where $-\infty \le a < b \le \infty$ and X is a continuous random variable as above.

That is, Y has the same distribution as X except that it is **truncated** on the left at a and on the right at b. Unlike X, Y cannot take values less than a or larger than b.

a. (3 marks) Mathematically show that the random variable W defined

$$W = Q_X(F_X(a) + U \times (F_X(b) - F_X(a)))$$

where U is a uniform random variable $U \sim U(0,1)$ has the same distribution as Y.

We can apply the $F_X()$ to W and disscuss the three cases. Note that $U \in [0,1]$.

Case 1: w < a

$$Pr(F_X(a) + U \times (F_X(b) - F_X(a)) \leq F_X(w)) = Pr(U \times (F_X(b) - F_X(a)) \leq F_X(w) - F_X(a)) = 0$$

Case 2: $a \le w \le b$

$$Pr(F_X(a) + U \times (F_X(b) - F_X(a)) \leq F_X(w)) = Pr(U \leq \frac{F_X(y) - F_X(a)}{F_X(b) - F_X(a)})$$

Case 3: b < w

$$Pr(F_X(a) + U \times (F_X(b) - F_X(a)) \leq F_X(w)) = Pr(U \times (F_X(b) - F_X(a)) \leq F_X(w) - F_X(a)) = 1 \leq Pr(F_X(a) + U \times (F_X(b) - F_X(a))) \leq Pr(U \times$$

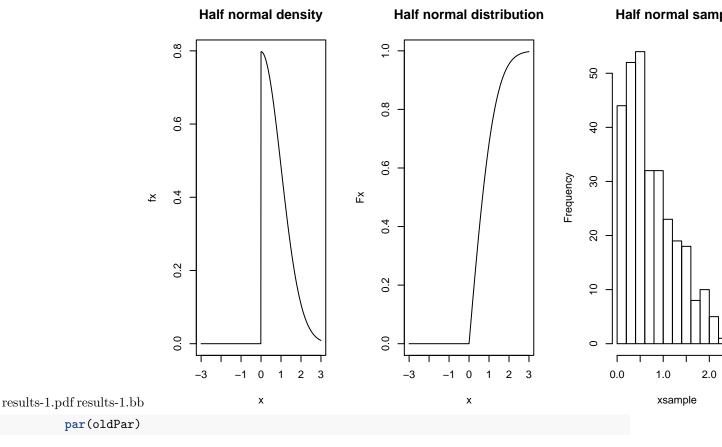
b. (15 marks) Here you are to write a function truncate() of the form

```
truncate <- function(ddist = dnorm, pdist = pnorm, qdist = qnorm, a = -Inf, b = Inf) {
    list(
        # density
        ddist = function(x) {
            den <- c()
            for (i in 1 : length(x)){</pre>
```

```
if (x[i] >= a & x[i] <= b){
          den[i] = (ddist(x[i]) / (pdist(b) - pdist(a)))
        } else {
          den[i] = 0
     return(den)
    },
    # distribution
    pdist = function(x, mean = 0,sd = 1) {
     distribution <- c()</pre>
     for (i in 1:length(x)){
        if (x[i] < a){</pre>
          distribution[i] = 0
        } else if (x[i] > b) {
         distribution[i] = 1
        }
        else {
          # calc by pdist(x)-pdist(a)]/[pdist(b)-pdist(a)]
          distribution[i] = (pdist(x[i],mean = mean,sd = sd) - pdist(a,mean = mean,sd = sd)) / (pdist
     }
     return(distribution)
    rdist = function(n,mean = 0, sd = 1) {
     return(qdist(pdist(a,mean = mean,sd = sd) + runif(n)*(pdist(b,mean = mean,sd = sd) - pdist(a,me
#list(ddist, pdist, rdist)
```

where `ddist`, `pdist`, and `qdist` refer to functions which calculate the density $f_X(x)$, dist. The function `truncate()` is to return a list with components named `ddist`, `pdist` and `rdist` c. That is, the following should work for the half-normal distribution.

```
half_normal <- truncate(a = 0)
xsample <- half_normal$rdist(300)
x <- seq(-3, 3, 0.01)
fx <- half_normal$ddist(x)
Fx <- half_normal$pdist(x)
oldPar <- par(mfrow = c(1,3))
plot(x, fx, type = "l", main = "Half normal density")
plot(x, Fx, type = "l", main = "Half normal distribution")
hist(xsample, main = "Half normal sample")</pre>
```

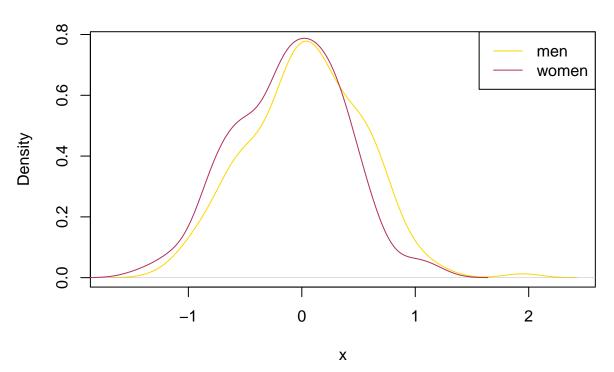


Hand in the above plots with your code.

- c. A 2011 article by Gil Greengross and Geoffrey Miller of the University of New Mexico was entitled "In the tested the sense of humour of 400 university students (200 men, 200 women) using a standardized The distributions of humour between men and women seems to be significantly different but what does
 - i. *(3 marks)* On a single (nicely labelled with a legend) draw the densities (in different colour

```
men <- rnorm(200, mean = 0.09, sd = 0.49)
women <- rnorm(200, mean = -0.09, sd = 0.49)
density_men = density(men)
density_women = density(women)
plot(density_men, col = "gold", xlab = 'x', main = "Densities for men and women")
lines(density_women, col = "maroon")
legend("topright", legend = c("men", "women"), col = c("gold", "maroon"), lwd = c(1,1))</pre>
```

Densities for men and women



- ii. *(4 marks)* Generate a random sample of 1000 scores from each of these distributions and save
 - the average humour ability of the men

x = rnorm(1000, mean = -0.09, sd = 0.49)

- the average humour ability of the women

Based on your sample, estimate the following

- the probability that the man will be funnier than the woman (at least as measured by this sc

```
y = rnorm(1000, mean = 0.09, sd =0.49)
results <- data.frame(women = x, men = y)

print("- the average humour ability of the men")

## [1] "- the average humour ability of the men"

print(mean(men))

## [1] 0.0387097

print("- the average humour ability of the women")</pre>
## [1] "- the average humour ability of the women"
```

[1] "- the average humour ability of the women"

print(mean(women))

[1] -0.1060419

print('- the probability that the man will be funnier than the woman (at least as measured by this scal

[1] "- the probability that the man will be funnier than the woman (at least as measured by this sca
count one by one....
counter = 0

```
for(i in 1:1000){
  if(y[i] > x[i]){
  counter <- counter + 1</pre>
}
prob <- counter / 1000
print(prob)
## [1] 0.61
     iii. *(4 marks)* [Are women funny?](https://www.theguardian.com/commentisfree/2014/mar/04/are-wom
          Generate 1000 pseudo random scores 'y' from the truncated distribution for men and another 10
             - the average humour ability of the men
             - the average humour ability of the women
             - the probability that the man will be funnier than the woman (at least as measured by thi
      funny <- truncate(a = 1.07)</pre>
      men <- funny$rdist(1000, mean = 0.09, sd = 0.49) #men
      women <- funny$rdist(1000, mean = -0.09, sd = 0.49) #women=
      res <- data.frame(women = women, men = men)
      mean_man <- mean(men)</pre>
      mean_women <- mean(women)</pre>
      count <- 0
      for(i in 1:1000){
        if(men[i] > women[i]){
        count <- count + 1</pre>
        }
      prob <- count/1000</pre>
      print('- the average humour ability of the men')
## [1] "- the average humour ability of the men"
     print(mean_man)
## [1] 1.254786
      print('- the average humour ability of the women')
## [1] "- the average humour ability of the women"
      print(mean_women)
## [1] 1.232599
      print('- the probability that the man will be funnier than the woman (at least as measured by thi
## [1] "- the probability that the man will be funnier than the woman (at least as measured by this sca
      print(prob)
## [1] 0.528
    iv. *(2 marks)* What conclusions do you draw about the differences between the humour of men and wo
```

There is no significant difference between the humor of men and women.

v. *(4 marks)* Repeat part iii, again conditioning on considering only individuals with a "humour a

```
funny2 <- truncate(a = 1.07)</pre>
      men <- funny2$rdist(1000,mean = 0.09,sd = 0.49 * 1.1) #men
      women <- funny2$rdist(1000,mean = 0.09,sd = 0.49) #women
      result <- data.frame(women = men, men = men)
      mean_men <- mean(men)</pre>
      mean_women <- mean(women)</pre>
      count <- 0
      count <- 0
      for(i in 1:1000){
        if(men[i] > women[i]){
        count <- count + 1</pre>
        }
      prob <- count/1000</pre>
      print('- the average humour ability of the men')
## [1] "- the average humour ability of the men"
      print(mean_man)
## [1] 1.254786
      print('- the average humour ability of the women')
## [1] "- the average humour ability of the women"
     print(mean_women)
## [1] 1.253287
print('- the probability that the man will be funnier than the woman (at least as measured by thi
## [1] "- the probability that the man will be funnier than the woman (at least as measured by this sca
     print(prob)
## [1] 0.56
```